SEQUENCE LISTING

<110> CALIFORNIA INSTITUTE OF TECHNOLOGY
 GUO, Zijian
 DUNPHY, William

<120> CHECKPOINT-ACTIVATING OLIGONUCLEOTIDES

<130> CIT1350-1

<140> US 09/849,617

<141> 2001-05-04

<150> US 60/202,028

<151> 2000-05-04

<160> 14

<170> PatentIn version 3.0

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adattequea aquetecque auceautige cauactgtat tittateuga tgetgetage 1140 tqttcaqtac ctccatqaaa atqqqqtgat acatcgtgat ctgaagcctg aaaatgtgct 1200 gttqtcatcc actagtgaag aatqttgcat aaagataacg gattttggac agtcaaaaat 1260 tctgggtgaa acgtctttaa tgagaacttt gtgtggaact cctacatact tggcgcctga 1320 agttttgaat acagcaggca caactggata cagtagtgca gtggattgct ggagtttagg 1380 agtcatcett tttgtgtgtc tttgtggata tccccccttt tcagaacaaa atagtaacat 1440 tecettgaaa aateagattg cagagggaaa atacacetae attgetgetg ettggagaaa 1500 tqtatcagaa caagcatttg atttagtcaa gaatcttctt gttgttgatc ctgagcaaag 1560 acttaccact aaacaagcac tqqaacatcc ctggcttcag gacgattcta tgaagcatac 1620 tgttgaaagg ttaatgtatg gggttgacca cacaatgcct cctccaatca agaaaaacat 1680 aattcgaaaa cggggacatg aatgggatca agatgccagt acttcatctt gctcagagat 1740 attaccaaca tcagccqaaa aqaqagcaaa aagataaaac aaaaaaaata cattgcgctt 1800 1845

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Ser Ser Gly Thr Leu Ser Ser Leu Asp Thr Val Pro Val Gln Asp Leu 35 40 45

Ala Ser Ile Pro Glu Asp Pro Glu Ile Asp Glu Asp Ile Pro Gln Pro 50 55 60

Trp Gly Arg Leu Trp Ala Leu Gly Lys Gly Phe Leu Asn His Asp Cys 70 75 80

Leu His Glu Glu Tyr Val Phe Gly Arg Asp Lys Lys Cys Asp Tyr Thr
85 90 95

Phe Asp Ile Pro Val Leu Asn Gln Thr Asp Arg Tyr Lys Thr Tyr Ser

Lys Arg His Phe Arg Ile Phe Gln Glu Leu Gly His Gly His Ser Arg 115 120 125

Val Ala Asn Ile Glu Asp Leu Ser Gly Asn Gly Thr Phe Val Asn Lys 130 135 140 Glu Ile Ile Gly Lys Gly Arg Thr Leu Pro Leu Thr Asn Asn Ala Glu 145 150 155 160

Ile Ala Leu Ser Leu Pro Thr Asn Lys Val Phe Val Phe Ser Asp Leu 165 170 175

Ser Val Asp Asp Gln Thr Ile Tyr Pro Lys Asp Phe Ile Asp Lys Tyr 180 185 190

Ile Met Ser Arg Pro Ile Gly Ser Gly Ala Cys Gly Glu Val Lys Leu 195 200 205

Ala Phe Gln Lys Ser Val Cys Lys Val Ala Val Lys Ile Ile Ser 210 215 220

Lys Arg Lys Phe Lys Met Asn Thr Ser Ser Asn Glu His Pro Ile Ser 225 230 235 240

Val Asp Thr Glu Ile Glu Ile Leu Lys Lys Leu Asp His Pro Cys Ile 245 250 255

Ile Lys Ile Glu Asn Phe Phe Asp Ser Glu Asp Phe Tyr Tyr Ile Val 260 265 270

Leu Glu Leu Met Glu Gly Gly Glu Leu Phe Asp Arg Val Val Asn Ser 275 280 285

Thr Arg Leu Arg Glu Pro Ile Ala Lys Leu Tyr Phe Tyr Gln Met Leu 290 295 300

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Lys Pro Glu Asn Val Leu Leu Ser Ser Thr Ser Glu Glu Cys Cys Ile 325 330 335

Lys Ile Thr Asp Phe Gly Gln Ser Lys Ile Leu Gly Glu Thr Ser Leu 340 345 350

Met Arg Thr Leu Cys Gly Thr Pro Thr Tyr Leu Ala Pro Glu Val Leu 355 360 365

Asn Thr Ala Gly Thr Thr Gly Tyr Ser Ser Ala Val Asp Cys Trp Ser 370 380

Leu Gly Val Ile Leu Phe Val Cys Leu Cys Gly Tyr Pro Pro Phe Ser 385 390 395 400

Glu Gln Asn Ser Asn Ile Pro Leu Lys Asn Gln Ile Ala Glu Gly Lys 405 410 415

Tyr Thr Tyr Ile Ala Ala Ala Trp Arg Asn Val Ser Glu Gln Ala Phe
420 425 430

Asp Leu Val Lys Asn Leu Leu Val Val Asp Pro Glu Gln Arg Leu Thr 435 440 445

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His Thr Val Glu Arg Leu Met Tyr Gly Val Asp His Thr Met Pro Pro

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475
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465
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Pro Ile Lys Lys Asn Ile Ile Arg Lys Arg Gly His Glu Trp Asp Gln
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      11
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gattatagta gagtagtaga gastattaga t		